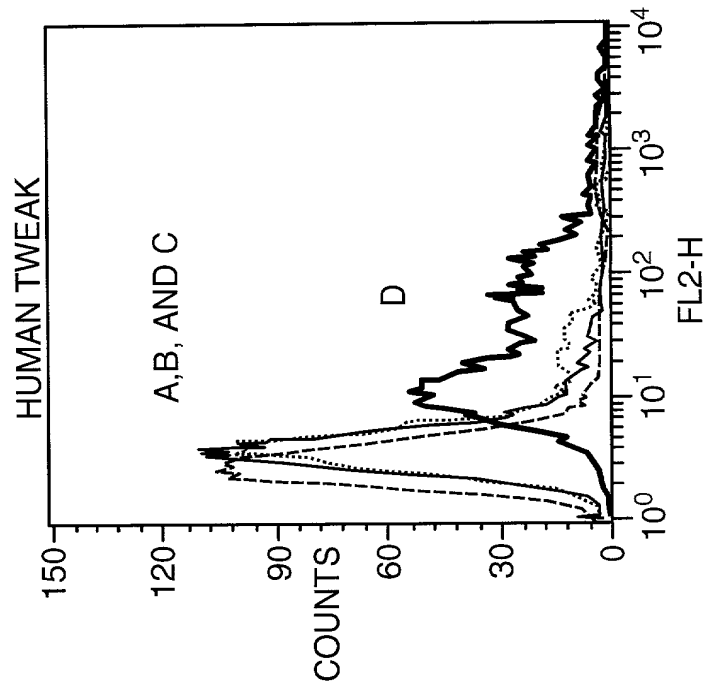
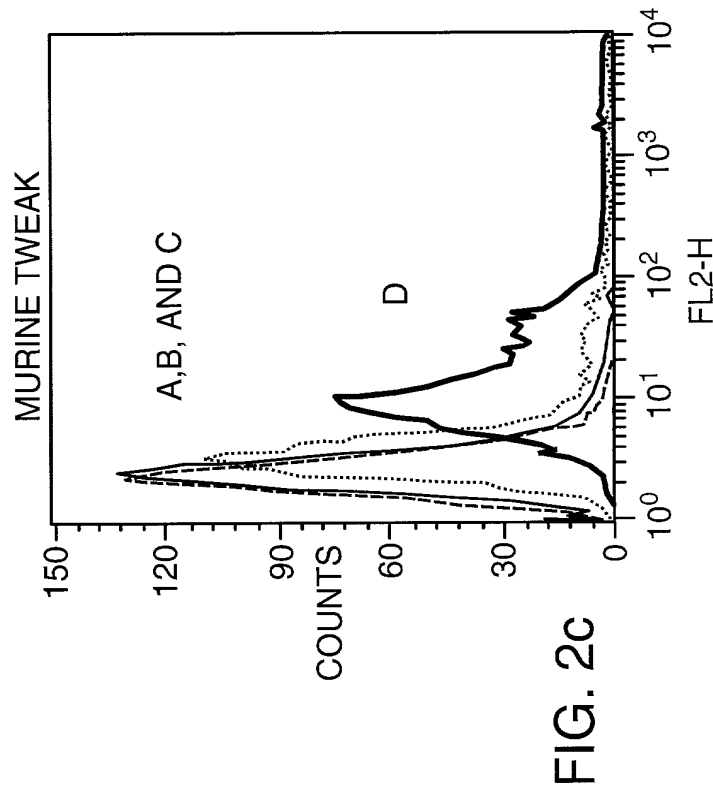
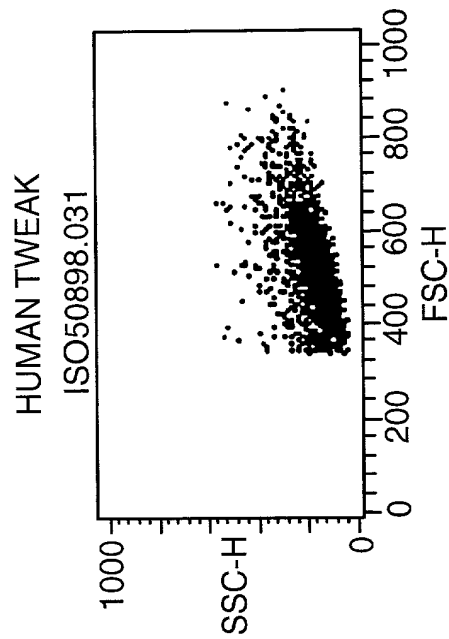
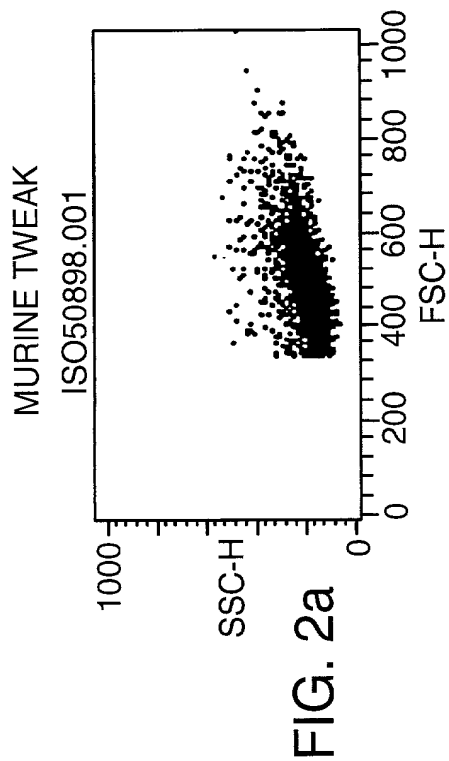


Human	1	MAARRSQRRRRGRRGPGTALLVPLALGLGLALACLGLLAVVSLGSRASL
Mouse	1	-----VLSGLALACLGLLVVVSLGSWATL
Human	51	SAQEPAQEELVAEEDQDPSELNPQTEESQDPAPFLNRLVRRPRRSAPKGRK
Mouse	27	SAQEPSQEELTAEDRRREPPELNPQTEESQDVVPFLEQLVRRPRRSAPKGRK
Human	101	TRARRAIAAHYEVHPRPGQDGAQAAGVDGTVSGWEEARINSSSPLRYNRQI
Mouse	77	ARPRRAIAAHYEVHPRPGQDGAQAAGVDGTVSGWEEKINSSSPLRYDRQI
Human	151	GEFIVTRAGLYLYLCQVHFDEGKAVYVKLDLLVDGVLAALRCLEEFSAATAA
Mouse	127	GEFTVIRAGLYLYLCQVHFDEGKAVYVKLDLLVNGVLAALRCLEEFSAATAA
Human	201	SSLGPQLRLCQVSGLLALRPGSSLRIRTLPWAHLKAAPFLTYFGLFQVH
Mouse	177	SSPGPQLRLCQVSGLLPLRPGSSLRIRTLPWAHLKAAPFLTYFGLFQVH

FIG. 1



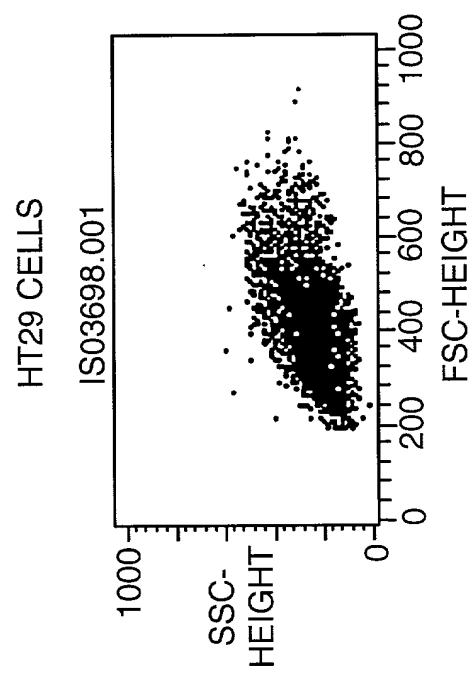


FIG. 3a

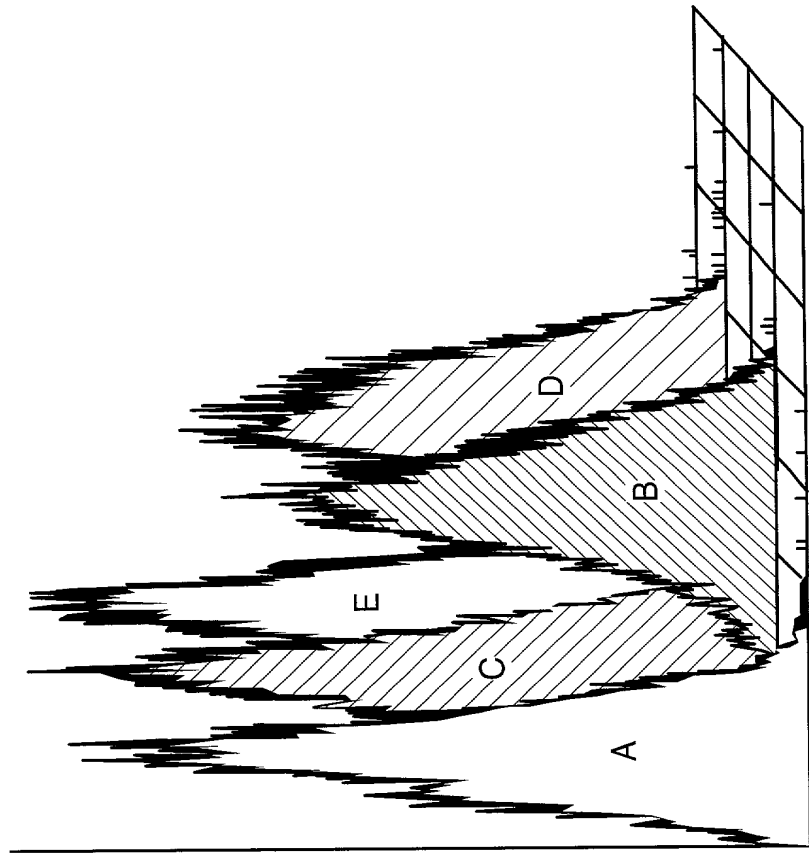


FIG. 3b

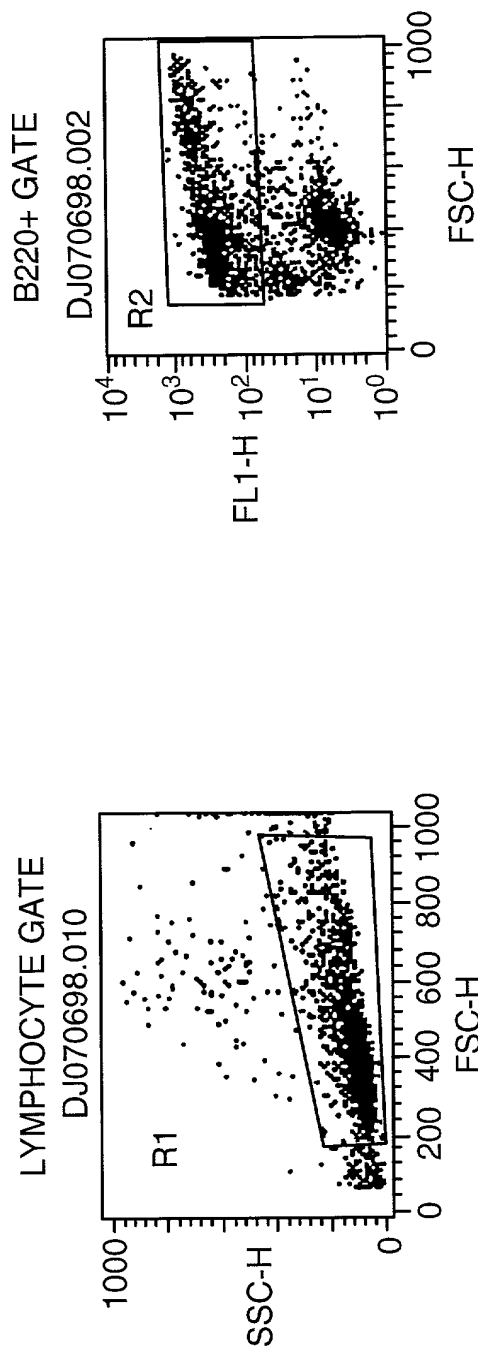


FIG. 4a

FIG. 4b

F1>F1 CONTROL

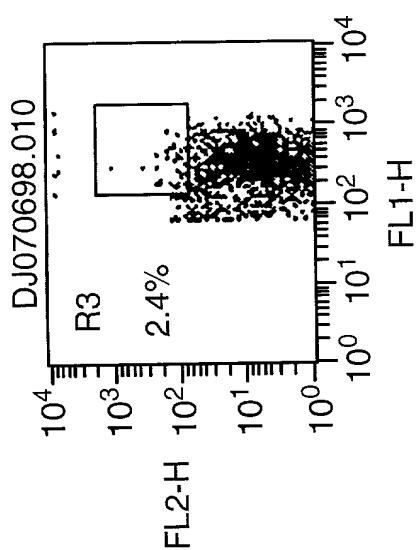


FIG. 4c

DBA/2>F1, UNTREATED

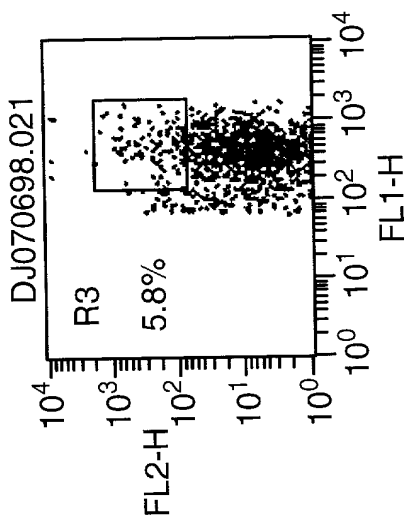


FIG. 4d

DBA/2>F1, Ha4/8

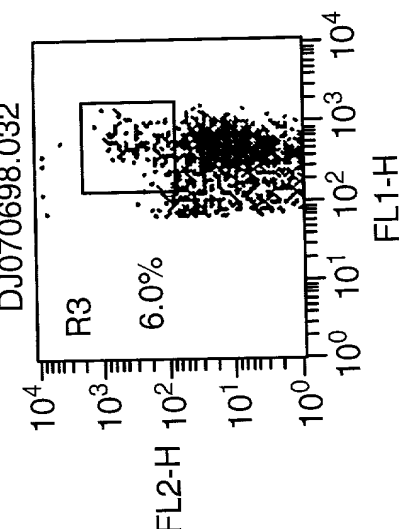


FIG. 4e

DBA/2>F1, MR1

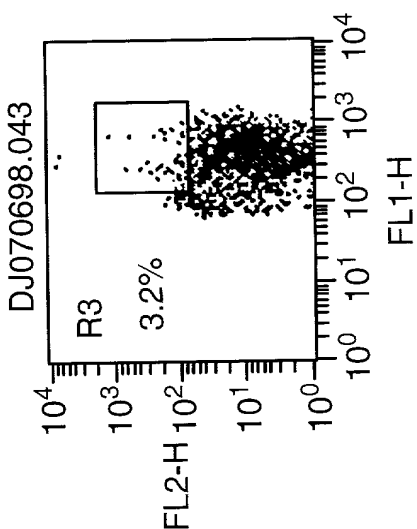


FIG. 4f

DBA/2>F1, AB.D3

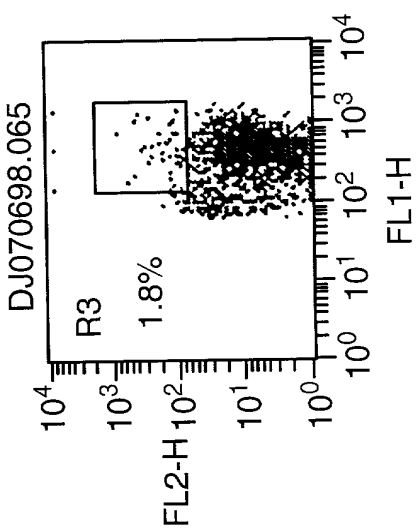


FIG. 4g